

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/591,726
Source: IFWP
Date Processed by STIC: 9/13/06

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 09/13/2006

PATENT APPLICATION: US/10/591,726

TIME: 11:07:38

Input Set : A:\21085064U1.TXT

Output Set: N:\CRF4\09132006\J591726.raw

4 <110> APPLICANT: UAB Research Foundation
 6 <120> TITLE OF INVENTION: BRHF1 AS A CANCER DIAGNOSTIC MARKER
 9 <130> FILE REFERENCE: 21085.0064P1
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/591,726
 C--> 11 <141> CURRENT FILING DATE: 2006-09-05
 11 <150> PRIOR APPLICATION NUMBER: 60/550,224
 12 <151> PRIOR FILING DATE: 2004-03-04
 14 <160> NUMBER OF SEQ ID NOS: 21
 16 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 18 <210> SEQ ID NO: 1
 19 <211> LENGTH: 105
 20 <212> TYPE: PRT
 21 <213> ORGANISM: Artificial Sequence
 23 <220> FEATURE:
 24 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
 25 Synthetic Construct
 27 <400> SEQUENCE: 1
 28 Met Lys Gly Leu Ser Pro Ile Ala Lys Gly Arg Lys Thr Ser Val Ser
 29 1 5 10 15
 30 Ala Ala Val Leu Val Ser Thr Thr Ile Pro Ile Ser Ser Val Trp Gly
 31 20 25 30
 32 Pro Leu Gln Ile Leu Gly Gln Lys Arg Gly Gln Lys Met Glu Gln Ala
 33 35 40 45
 34 Asn His Pro Val Gly Leu Asp Ile Ser Val Val Tyr Lys Asp Thr Leu
 35 50 55 60
 36 Lys Lys Ile Val Gln Gln Glu Thr Ser Cys Pro Phe Thr His Val His
 37 65 70 75 80
 38 Tyr Ala Glu Gly Ile Thr Gly Arg His Thr Ala Pro Glu Asp Glu Gly
 39 85 90 95
 40 Ser Leu Ala Gln Lys Pro Pro Ile Arg
 41 100 105
 44 <210> SEQ ID NO: 2
 45 <211> LENGTH: 270
 46 <212> TYPE: PRT
 47 <213> ORGANISM: Artificial Sequence
 49 <220> FEATURE:
 50 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
 51 Synthetic Construct
 53 <400> SEQUENCE: 2
 54 Met Asn Ile Asp Ala Lys Ile Leu Asn Lys Ile Leu Ala Asn Gln Ile
 55 1 5 10 15
 56 Gln Gln His Ile Lys Lys Leu Ile His His Asp Gln Val Gly Phe Ile
 57 20 25 30

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58 Pro Gly Met Gln Gly Trp Phe Asn Ile His Lys Ser Ile Asn Val Ile
59          35          40          45
60 Gln His Ile Asn Arg Thr Lys Asp Lys Asn His Met Ile Ile Ser Val
61          50          55          60
62 Asp Ala Glu Lys Ala Phe Asp Lys Val Gln Gln His Phe Met Leu Lys
63 65          70          75          80
64 Thr Leu Asn Lys Leu Gly Ile Asp Gly Thr Tyr Leu Lys Ile Ile Arg
65          85          90          95
66 Ala Ile Tyr Asp Lys Pro Thr Ala Asn Ile Ile Leu Asn Gly Leu Lys
67          100          105          110
68 Leu Glu Ala Phe Pro Leu Lys Thr Gly Thr Arg Gln Gly Cys Pro Leu
69          115          120          125
70 Ser Leu Leu Leu Phe Asn Ile Val Leu Glu Val Leu Ala Arg Ala Ile
71          130          135          140
72 Arg Gln Glu Lys Glu Ile Asn Cys Ile Gln Leu Gly Lys Glu Glu Val
73 145          150          155          160
74 Lys Leu Pro Leu Phe Ala Asp Asp Met Ile Val Tyr Leu Glu Asn Pro
75          165          170          175
76 Val Val Ser Ala Pro Asn Leu Leu Lys Leu Ile Ser Asn Phe Ser Lys
77          180          185          190
78 Val Ser Gly Tyr Lys Ile Asn Val Gln Lys Ser Gln Ala Phe Leu Tyr
79          195          200          205
80 Thr Asn Asn Arg Gln Thr Glu Ser Gln Ile Met Ser Glu Leu Pro Phe
81          210          215          220
82 Thr Ile Ala Ser Lys Arg Ile Lys Tyr Leu Gly Ile Gln Leu Thr Arg
83 225          230          235          240
84 Asp Val Lys Asp Leu Phe Lys Glu Asn Tyr Lys Pro Leu Leu Asn Glu
85          245          250          255
86 Ile Lys Glu Asp Thr Asn Lys Cys Lys Asn Ile Pro Cys Ser
87          260          265          270
90 <210> SEQ ID NO: 3
91 <211> LENGTH: 315
92 <212> TYPE: DNA
93 <213> ORGANISM: Artificial Sequence
95 <220> FEATURE:
96 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
97     Synthetic Construct
99 <400> SEQUENCE: 3
100 atgaagggat tatcgccctat cgccaagggg aggaaaacta gtgtttctgc tgctgtgttg 60
101 gtgagcacia ctattccgat cagcagtgtc tggggaccat tgcagattct tgggcaaaag 120
102 agaggacaga aaatggagca ggccaatcac ccagtggggc ttgatatcag tgtggtttac 180
103 aaggacacct taaaaaagat tgtccaacaa gaaacaagct gccccttcac ccatgtccac 240
104 tatgctgagg gaatcactgg aaggcacact gcccagagg atgaaggttc tctggcccag 300
105 aagcccccaa tcaga 315
107 <210> SEQ ID NO: 4
108 <211> LENGTH: 810
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence
112 <220> FEATURE:

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113 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =

114 Synthetic Construct

116 <400> SEQUENCE: 4

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117 atgaacatcg atgcaaaaat cctcaataaa atactggcaa accaaatcca gcagcacatc 60
118 aaaaagctta tccaccatga tcaagtgggc ttcattccctg ggatgcaagg ctgggttcaac 120
119 atacacaaat caataaatgt aatccagcat ataaacagaa ccaaagacaa aaaccacatg 180
120 attatctcag tagatgcaga aaaggccttt gacaaagtgc aacaacactt catgctaaaa 240
121 actctcaata aattaggtat tgatgggacg tatctcaaaa taataagagc tatctatgac 300
122 aaaccacacag ccaatatcat actgaatggg ctaaaactgg aagcattccc ttgaaaact 360
123 ggcacaagac agggatgccc tctctcactt ctctattca acatagtgtt ggaagtctctg 420
124 gccagggcaa tcaggcagga gaaggaaata aattgtattc aattaggaaa agaggaagtt 480
125 aaattgcccc tgtttgacga tgacatgatt gtatatctgg aaaaccccgt cgtctcagcc 540
126 ccaaatctcc ttaagctgat aagcaacttc agcaaagtct caggatacaa aatcaacgtg 600
127 caaaaatcac aagcattctt atacaccaat aacagacaaa cagagagcca aatcatgagt 660
128 gaactcccat tcacaattgc ttcaaagaga ataaaatacc taggaatcca acttacaagg 720
129 gatgtgaagg acctcttcaa ggagaactac aaaccactgc tcaacgaaat aaaagaggat 780
130 acaaacaaat gcaagaacat tccatgctca 810

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132 <210> SEQ ID NO: 5

133 <211> LENGTH: 1263

134 <212> TYPE: DNA

135 <213> ORGANISM: Artificial Sequence

137 <220> FEATURE:

138 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =

139 Synthetic Construct

141 <400> SEQUENCE: 5

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142 catctacaga actctccacc ccaaatacaac agaataataca ttttttttcag caccacacca 60
143 cacctattcc aaaattgacc acatagtgtg aagtaaagct ctctcagca aatgtaaaag 120
144 aacagaaatt ataacaaact atctctcaga ccacagtgc atcaaaactag aactcaggat 180
145 taagaatctc actcaaagcc gctcaactac atggaaactg aacaacctgc tcctgaatga 240
146 ctactgggta cataacgaaa tgaaggcaga aataaagatg ttctttgaaa ccaacgagaa 300
147 caaagacacc acataccaga atctctggga cgcattcaaa gcagtgtgta gagggaaatt 360
148 tatagcacta aatgcctacc agagaaagca ggaaagatcc aaaattgaca ccctaacatc 420
149 acaattaaaa gaactagaaa agcaagagca aacacattca aaagctagca gaaggcaaga 480
150 aataactaaa atcagagcag aactgaagga aatagagaca caaaaaaccc ttcaaaaaat 540
151 caatgaatcc aggagctggt tttttgaaag gatcaacaaa attgatagac cgctagcaag 600
152 actaataaag aaaaaaagag agaagaatca aatagacaca ataaaaaatg ataaagggga 660
153 tatcaccacc gatcccacag aaatacaaac taccatcaga gaatactaca aacacctcta 720
154 cgcaaataaa ctagaaaatc tggaagaaat ggatacattc ctgcacacat acactctccc 780
155 aagactaaac caggaagaag ttgaatctct gaatcgacca ataacaggct ctgaaattgt 840
156 ggcaataatc aatagtttac caacccaaaa gagtccagga ccagatggat tcacagccga 900
157 attctaccag aggtacaagg aggaactggt accattcctt ctgaaactat tccaatcaat 960
158 agaaaaagag ggaatcctcc ctaactcatt ttatgagacc agcatcattc tgataccaaa 1020
159 gccgggcaga gacacaacca aaaaagagaa ttttagacca atatccttga tgaacattga 1080
160 tgcaaaaatc ctcaataaaa tactggcaaa ccgaatccag cagcacatca aaaagcttat 1140
161 ccaccatgat caagtgggct tcatccctgg gatgcaaggc tgggttcaata tacgcaaadc 1200
162 aataaatgta atccagcata taaacagagc caaagacaaa aaccacatga ttatctcaat 1260
163 aga 1263

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165 <210> SEQ ID NO: 6

166 <211> LENGTH: 10

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167 <212> TYPE: DNA
168 <213> ORGANISM: Artificial Sequence
170 <220> FEATURE:
171 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
172     Synthetic Construct
174 <400> SEQUENCE: 6
175 cagagcctgt                                     10
177 <210> SEQ ID NO: 7
178 <211> LENGTH: 10
179 <212> TYPE: DNA
180 <213> ORGANISM: Artificial Sequence
182 <220> FEATURE:
183 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
184     Synthetic Construct
186 <400> SEQUENCE: 7
187 ctctgggaca                                     10
189 <210> SEQ ID NO: 8
190 <211> LENGTH: 375
191 <212> TYPE: PRT
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
196     Synthetic Construct
198 <400> SEQUENCE: 8
199 Met  Lys  Gly  Leu  Ser  Pro  Ile  Ala  Lys  Gly  Arg  Lys  Thr  Ser  Val  Ser
200   1      5      10      15
201 Ala  Ala  Val  Leu  Val  Ser  Thr  Thr  Ile  Pro  Ile  Ser  Ser  Val  Trp  Gly
202      20      25      30
203 Pro  Leu  Gln  Ile  Leu  Gly  Gln  Lys  Arg  Gly  Gln  Lys  Met  Glu  Gln  Ala
204      35      40      45
205 Asn  His  Pro  Val  Gly  Leu  Asp  Ile  Ser  Val  Val  Tyr  Lys  Asp  Thr  Leu
206      50      55      60
207 Lys  Lys  Ile  Val  Gln  Gln  Glu  Thr  Ser  Cys  Pro  Phe  Thr  His  Val  His
208      65      70      75      80
209 Tyr  Ala  Glu  Gly  Ile  Thr  Gly  Arg  His  Thr  Ala  Pro  Glu  Asp  Glu  Gly
210      85      90      95
211 Ser  Leu  Ala  Gln  Lys  Pro  Pro  Ile  Arg  Met  Asn  Ile  Asp  Ala  Lys  Ile
212      100     105     110
213 Leu  Asn  Lys  Ile  Leu  Ala  Asn  Gln  Ile  Gln  Gln  His  Ile  Lys  Lys  Leu
214      115     120     125
215 Ile  His  His  Asp  Gln  Val  Gly  Phe  Ile  Pro  Gly  Met  Gln  Gly  Trp  Phe
216      130     135     140
217 Asn  Ile  His  Lys  Ser  Ile  Asn  Val  Ile  Gln  His  Ile  Asn  Arg  Thr  Lys
218      145     150     155     160
219 Asp  Lys  Asn  His  Met  Ile  Ile  Ser  Val  Asp  Ala  Glu  Lys  Ala  Phe  Asp
220      165     170     175
221 Lys  Val  Gln  Gln  His  Phe  Met  Leu  Lys  Thr  Leu  Asn  Lys  Leu  Gly  Ile
222      180     185     190
223 Asp  Gly  Thr  Tyr  Leu  Lys  Ile  Ile  Arg  Ala  Ile  Tyr  Asp  Lys  Pro  Thr

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224	195	200	205
225	Ala Asn Ile Ile Leu Asn Gly Leu Lys Leu Glu Ala Phe Pro Leu Lys		
226	210	215	220
227	Thr Gly Thr Arg Gln Gly Cys Pro Leu Ser Leu Leu Leu Phe Asn Ile		
228	225	230	235
229	Val Leu Glu Val Leu Ala Arg Ala Ile Arg Gln Glu Lys Glu Ile Asn		
230	245	250	255
231	Cys Ile Gln Leu Gly Lys Glu Glu Val Lys Leu Pro Leu Phe Ala Asp		
232	260	265	270
233	Asp Met Ile Val Tyr Leu Glu Asn Pro Val Val Ser Ala Pro Asn Leu		
234	275	280	285
235	Leu Lys Leu Ile Ser Asn Phe Ser Lys Val Ser Gly Tyr Lys Ile Asn		
236	290	295	300
237	Val Gln Lys Ser Gln Ala Phe Leu Tyr Thr Asn Asn Arg Gln Thr Glu		
238	305	310	315
239	Ser Gln Ile Met Ser Glu Leu Pro Phe Thr Ile Ala Ser Lys Arg Ile		
240	325	330	335
241	Lys Tyr Leu Gly Ile Gln Leu Thr Arg Asp Val Lys Asp Leu Phe Lys		
242	340	345	350
243	Glu Asn Tyr Lys Pro Leu Leu Asn Glu Ile Lys Glu Asp Thr Asn Lys		
244	355	360	365
245	Cys Lys Asn Ile Pro Cys Ser		
246	370	375	
249	<210> SEQ ID NO: 9		
250	<211> LENGTH: 1125		
251	<212> TYPE: DNA		
252	<213> ORGANISM: Artificial Sequence		
254	<220> FEATURE:		
255	<223> OTHER INFORMATION: Description of Artificial Sequence:/note =		
256	Synthetic Construct		
258	<400> SEQUENCE: 9		
259	atgaagggat tatcgcttat cgccaagggg aggaaaacta gtgtttctgc tgctgtgttg	60	
260	gtgagcacia ctattccgat cagcagtgtc tggggaccat tgcagattct tgggcaaaag	120	
261	agaggacaga aaatggagca ggccaatcac ccagtggggc ttgatatacag tgtggtttac	180	
262	aaggacacct taaaaaagat tgtccaacaa gaaacaagct gccccttcac ccatgtccac	240	
263	tatgctgagg gaatcactgg aaggcacact gcccagagg atgaagggtt tctggcccag	300	
264	aagcccccaa tcagaatgaa catcgatgca aaaatcctca ataaaataact ggcaaaccac	360	
265	atccagcagc acatcaaaaa gcttatccac catgatcaag tgggcttcat ccctgggatg	420	
266	caaggctggt tcaacataca caaatcaata aatgtaatcc agcatataaa cagaacacaa	480	
267	gacaaaaacc acatgattat ctcatgatg gcagaaaagg cctttgacaa agttcaacaa	540	
268	cacttcatgc taaaaactct caataaatta ggtattgatg ggacgtatct caaaataata	600	
269	agagctatct atgacaaacc cacagccaat atcatactga atgggctaaa actggaagca	660	
270	ttccctttga aaactggcac aagacaggga tgccctctct cacttctcct attcaacata	720	
271	gtgttggaag ttctggccag ggcaatcagg caggagaagg aaataaattg tattcaatta	780	
272	ggaaaagagg aagttaaatt gcccctgttt gcagatgaca tgattgtata tctggaaaac	840	
273	cccgctcgtct cagccccaaa tctccttaag ctgataagca acttcagcaa agtctcagga	900	
274	tacaaaatca acgtgcaaaa atcacaagca ttcttatata ccaataacag acaaacagag	960	
275	agccaaatca tgagtgaact ccatttcaca attgcttcaa agagaataaa atacctagga	1020	
276	atccaactta caagggatgt gaaggacctc ttcaaggaga actacaaacc actgctcaac	1080	

VERIFICATION SUMMARY

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L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date